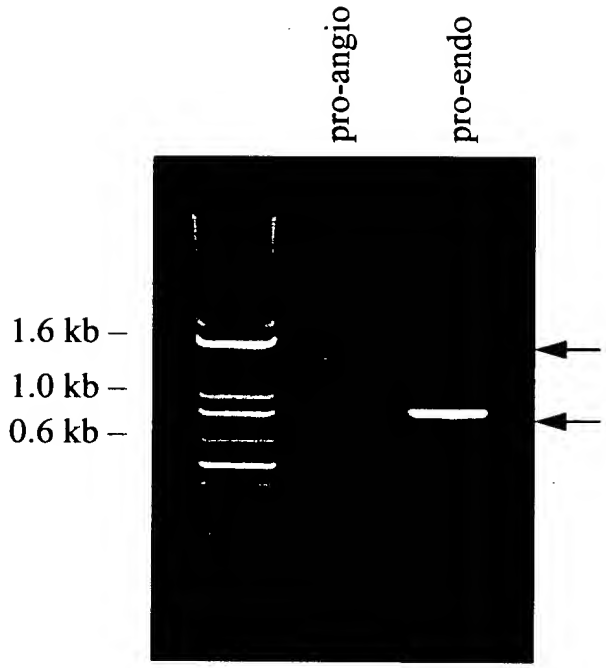


FIG. 1

```
1                                                                    80
CCCTGGCGGGCAGATGACATCCTGGCCGGCCCCCCGCGCCTGCTGGACCCCCAGCCCTACCCCGGGGCCCCGCACCACGG
81                                                                  160
CTCCTACGTGCACTTCCAGCCGGCTCGCCCCACTGGTGGGCCCGTCCACACCCACACCCACACCCACCAGGACTTCCAGC
161                                                                 240
TGGTGCTGCACCTGGTGGCCCTGAACAGCCCGCAGCCGGGCGGCATGCGAGGCATCCGGGGAGCGGACTTCCAGTGCTTC
241                                                                 320
CAGCAGGCGCGCGCCGCGGGGCTGGCCGGCACCTTCCGGGCCTTCCTGTCGTCGCGGCTGCAGGACCTCTACAGCATCGT
321                                                                 400
GCGCCGCGCCGACCGCACCGGGGTGCCCGTCGTCAACCTCAGGGACGAGGTGCTCTTCCCCAGCTGGGAGGCCTTATTCT
401                                                                 480
CGGGCTCCGAGGGCCAGCTGAAGCCCGGGGCCCGCATCTTCTCTTTCGACGGCAGAGATGTCCTGCAGCACCCCGCCTGG
481                                                                 560
CCCCGGAAGAGCGTGTGGCACGGCTCCGACCCCAGCGGGCGCCGCCTGACCGACAGCTACTGCGAGACGTGGCGGACGGA
561                                                                 640
GGCCCCGGCGGCCACCGGGCAGGCGTCGTCGCTGCTGGCGGGCAGGCTGCTGGAGCAGGAGGCCGCGAGCTGCCGCCACG
641                                                                 720
CCTTCGTGGTGCTCTGCATCGAGAACAGCGTCATGACCTCCTTCTCCAAGTAGGGCCGCGCGGCCCACGGACAGGCGGGG
721                                                                 800
GAGGGGGCGCCCGCAGGAGCATCCGCCGCCCCGGGGGGGCCTGGCCGGGACGCTTGCCTGCACCGTCACGTTTAATGTAA
801                                                                 829
TCCTCAAGAAATAAAAGGAAGCCAAAGAG
```

FIG. 2

```
   1 ccctggcgggcagatgacatcctggccggccccccgcgcctgctg
     P  W  R  A  D  D  I  L  A  G  P  P  R  L  L    15
  46 gaccccagccctaccccgggggccccgcaccacggctcctacgtg
     D  P  Q  P  Y  P  G  A  P  H  H  G  S  Y  V    30
  91 cacttccagccggctcgccccactggtgggcccgtccacacccac
     H  F  Q  P  A  R  P  T  G  G  P  V  H  T  H    45
 136 acccacacccaccaggacttccagctggtgctgcacctggtggcc
     T  H  T  H  Q  D  F  Q  L  V  L  H  L  V  A    60
 181 ctgaacagcccgcagccgggcggcatgcgaggcatccggggagcg
     L  N  S  P  Q  P  G  G  M  R  G  I  R  G  A    75
 226 gacttccagtgcttccagcaggcgcgccgccggggctggccggc
     D  F  Q  C  F  Q  Q  A  R  A  A  G  L  A  G    90
 271 accttccgggccttcctgtcgtcgcggctgcaggacctctacagc
     T  F  R  A  F  L  S  S  R  L  Q  D  L  Y  S   105
 316 atcgtgcgccgcgccgaccgcaccggggtgcccgtcgtcaacctc
     I  V  R  R  A  D  R  T  G  V  P  V  V  N  L   120
 361 agggacgaggtgctcttccccagctgggaggccttattctcgggc
     R  D  E  V  L  F  P  S  W  E  A  L  F  S  G   135
 406 tccgagggccagctgaagcccggggcccgcatcttctctttcgac
     S  E  G  Q  L  K  P  G  A  R  I  F  S  F  D   150
 451 ggcagagatgtcctgcagcaccccgcctggccccggaagagcgtg
     G  R  D  V  L  Q  H  P  A  W  P  R  K  S  V   165
 496 tggcacggctccgaccccagcgggcgccgcctgaccgacagctac
     W  H  G  S  D  P  S  G  R  R  L  T  D  S  Y   180
 541 tgcgagacgtggcggacggaggccccggcggccaccgggcaggcg
     C  E  T  W  R  T  E  A  P  A  A  T  G  Q  A   195
 586 tcgtcgctgctggcgggcaggctgctggagcaggaggccgcgagc
     S  S  L  L  A  G  R  L  L  E  Q  E  A  A  S   210
 631 tgccgccacgccttcgtggtgctctgcatcgagaacagcgtcatg
     C  R  H  A  F  V  V  L  C  I  E  N  S  V  M   225
 676 acctccttctccaagtagggccgcgcggcccacggacaggcgggg
     T  S  F  S  K  *                              230
 721 gaggggggcgcccgcaggagcatccgccgccccgggggggcctggc
 766 cgggacgcttgcctgcaccgtcacgtttaatgtaatcctcaagaa
 811 ataaaaggaagccaaagag
```

FIG. 3

```
1                                                                    80
CACACCCACCAGGACTTCCAGCTGGTGCTGCACCTGGTGGCCCTGAACAGCCCGCAGCCGGGCGGCATGCGAGGCATCCG
81                                                                   160
GGGAGCGGACTTCCAGTGCTTCCAGCAGGCGCGCGCCGCGGGGCTGGCCGGCACCTTCCGGGCCTTCCTGTCGTCGCGGC
161                                                                  240
TGCAGGACCTCTACAGCATCGTGCGCCGCGCCGACCGCACCGGGGTGCCCGTCGTCAACCTCAGGGACGAGGTGCTCTTC
241                                                                  320
CCCAGCTGGGAGGCCTTATTCTCGGGCTCCGAGGGCCAGCTGAAGCCCGGGGCCCGCATCTTCTCTTTCGACGGCAGAGA
321                                                                  400
TGTCCTGCAGCACCCCGCCTGGCCCCGGAAGAGCGTGTGGCACGGCTCCGACCCCAGCGGGCGCCGCCTGACCGACAGCT
401                                                                  480
ACTGCGAGACGTGGCGGACGGAGGCCCCGGCGGCCACCGGGCAGGCGTCGTCGCTGCTGGCGGGCAGGCTGCTGGAGCAG
481                                                                  555
GAGGCCGCGAGCTGCCGCCACGCCTTCGTGGTGCTCTGCATCGAGAACAGCGTCATGACCTCCTTCTCCAAGTAG
```

FIG. 4

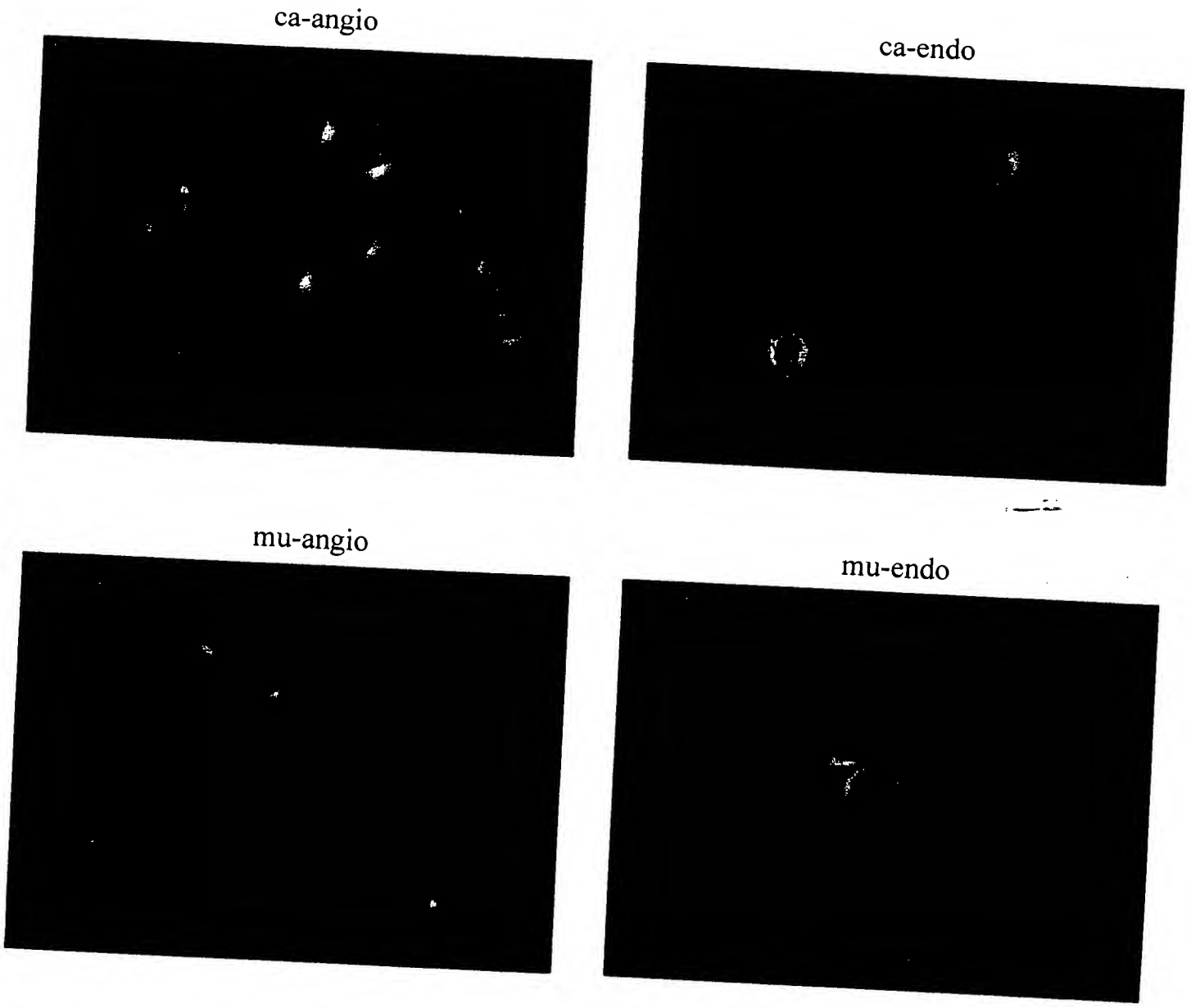
```
  1 cacacccaccaggacttccagctggtgctgcacctggtggccctg
    H  T  H  Q  D  F  Q  L  V  L  H  L  V  A  L   15
 46 aacagcccgcagccgggcggcatgcgaggcatccggggagcggac
    N  S  P  Q  P  G  G  M  R  G  I  R  G  A  D   30
 91 ttccagtgcttccagcaggcgcgcgccgcggggctggccggcacc
    F  Q  C  F  Q  Q  A  R  A  A  G  L  A  G  T   45
136 ttccgggccttcctgtcgtcgcggctgcaggacctctacagcatc
    F  R  A  F  L  S  S  R  L  Q  D  L  Y  S  I   60
181 gtgcgccgcgccgaccgcaccggggtgcccgtcgtcaacctcagg
    V  R  R  A  D  R  T  G  V  P  V  V  N  L  R   75
226 gacgaggtgctcttccccagctgggaggccttattctcgggctcc
    D  E  V  L  F  P  S  W  E  A  L  F  S  G  S   90
271 gagggccagctgaagcccggggcccgcatcttctctttcgacggc
    E  G  Q  L  K  P  G  A  R  I  F  S  F  D  G  105
316 agagatgtcctgcagcaccccgcctggccccggaagagcgtgtgg
    R  D  V  L  Q  H  P  A  W  P  R  K  S  V  W  120
361 cacggctccgaccccagcgggcgccgcctgaccgacagtactgc
    H  G  S  D  P  S  G  R  R  L  T  D  S  Y  C  135
406 gagacgtggcggacggaggccccggcggccaccgggcaggcgtcg
    E  T  W  R  T  E  A  P  A  A  T  G  Q  A  S  150
451 tcgctgctggcgggcaggctgctggagcaggaggccgcgagctgc
    S  L  L  A  G  R  L  L  E  Q  E  A  A  S  C  165
496 cgccacgccttcgtggtgctctgcatcgagaacagcgtcatgacc
    R  H  A  F  V  V  L  C  I  E  N  S  V  M  T  180
541 tccttctccaagtag
    S  F  S  K  *                                184
```

FIG. 5

```
1    HTHQDFQLVLHLVALNSPQPGGMRGIRGADFQCFQQARAAGLAGTFRAFLSSRLQDLYSI  endostatin-canine.PRO
1    HVHQDFQPALHLVALNTPLSGGMRGIRGADFQCFQQARQVGLAGTFRAFLSSRLQDLYSI  endostatin-chicken.PRO
1    HSHRDFQPVLHLVALNSPLSGGMRGIRGADFQCFQQARAVGLAGTFRAFLSSRLQDLYSI  endostatin-human.PRO
1    HTHQDFQPVLHLVALNTPLSGGMRGIRGADFQCFQQARAVGLSGTFRAFLSSRLQDLYSI  endostatin-mouse.PRO

61   VRRADRTGVPVVNLRDEVLFPSWEALFSGSEGQLKPGARIFSFDGRDVLQHPAWPRKSVW  endostatin-canine.PRO
61   VRRADRTAVPIVNLRDEVLFSNWEALFTGSEAPLRAGARILSFDGRDILQDSAWPQKSIW  endostatin-chicken.PRO
61   VRRADRAAVPIVNLKDELLFPSWEALFSGSEGPLKPGARIFSFDGKDVLRHPIWPQKSVW  endostatin-human.PRO
61   VRRADRGSVPIVNLKDEVLSPSWDSLFSGSQGQLQPGARIFSFDGRDVLRHPAWPQKSVW  endostatin-mouse.PRO

121  HGSDPSGRRLTDSYCETWRTEAPAATGQASSLLAGRLLEQEAASCRHAFVVLCIENSVMT  endostatin-canine.PRO
121  HGSDAKGRRLPESYCEAWRTDERGTSGQASSLSSGKLLEQSASSCQHAFVVLCIENSFMT  endostatin-chicken.PRO
121  HGSDPNGRRLTESYCETWRTEAPSATGQASSLLGGRLLGQSAASCHHAYIVLCIENSFMT  endostatin-human.PRO
121  HGSDPSGRRLMESYCETWRTETTGATGQASSLLSGRLLEQKAASCHNSYIVLCIENSFMT  endostatin-mouse.PRO

181  SFSK                                                           endostatin-canine.PRO
181  AAKK                                                           endostatin-chicken.PRO
181  ASK                                                            endostatin-human.PRO
181  SFSK                                                           endostatin-mouse.PRO
```

FIG. 6

FIG. 7



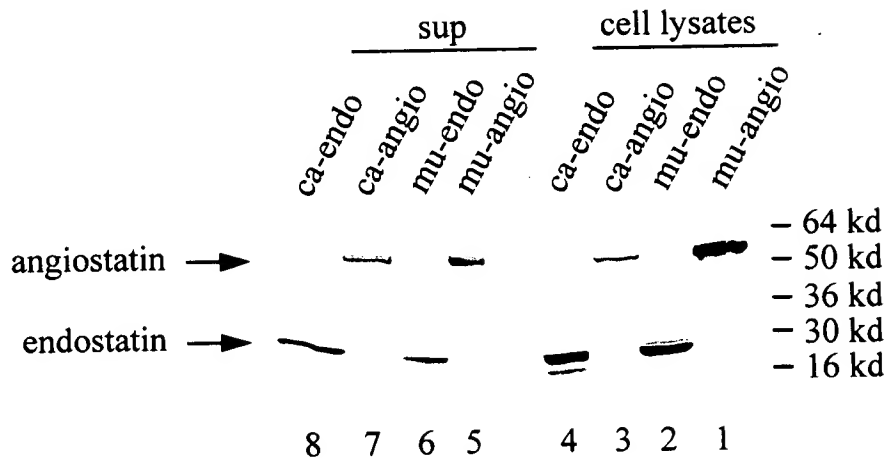

sup
cell lysates
ca-endo
ca-angio
mu-endo
mu-angio
ca-endo
ca-angio
mu-endo
mu-angio
angiostatin
endostatin
64 kd
50 kd
36 kd
30 kd
16 kd
8 7 6 5 4 3 2 1

FIG. 8

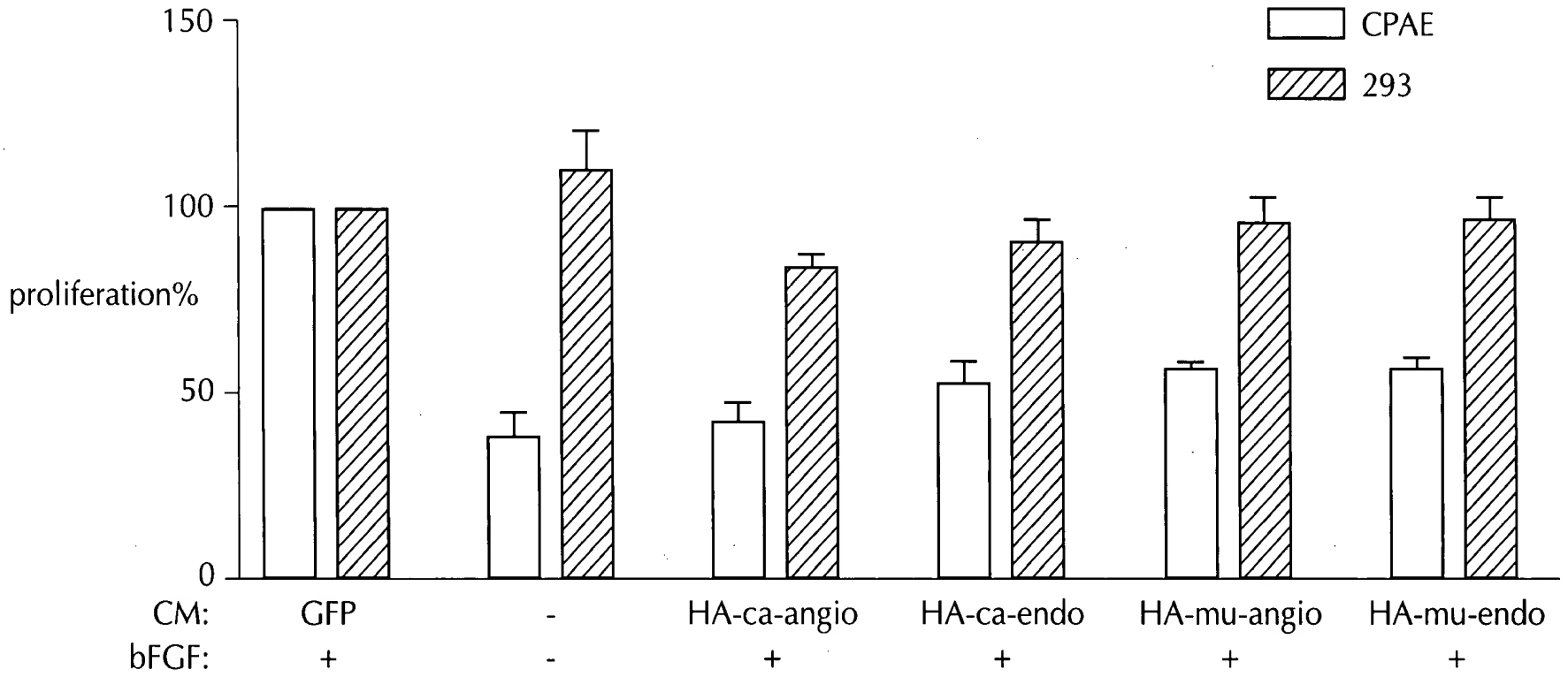


FIG. 9

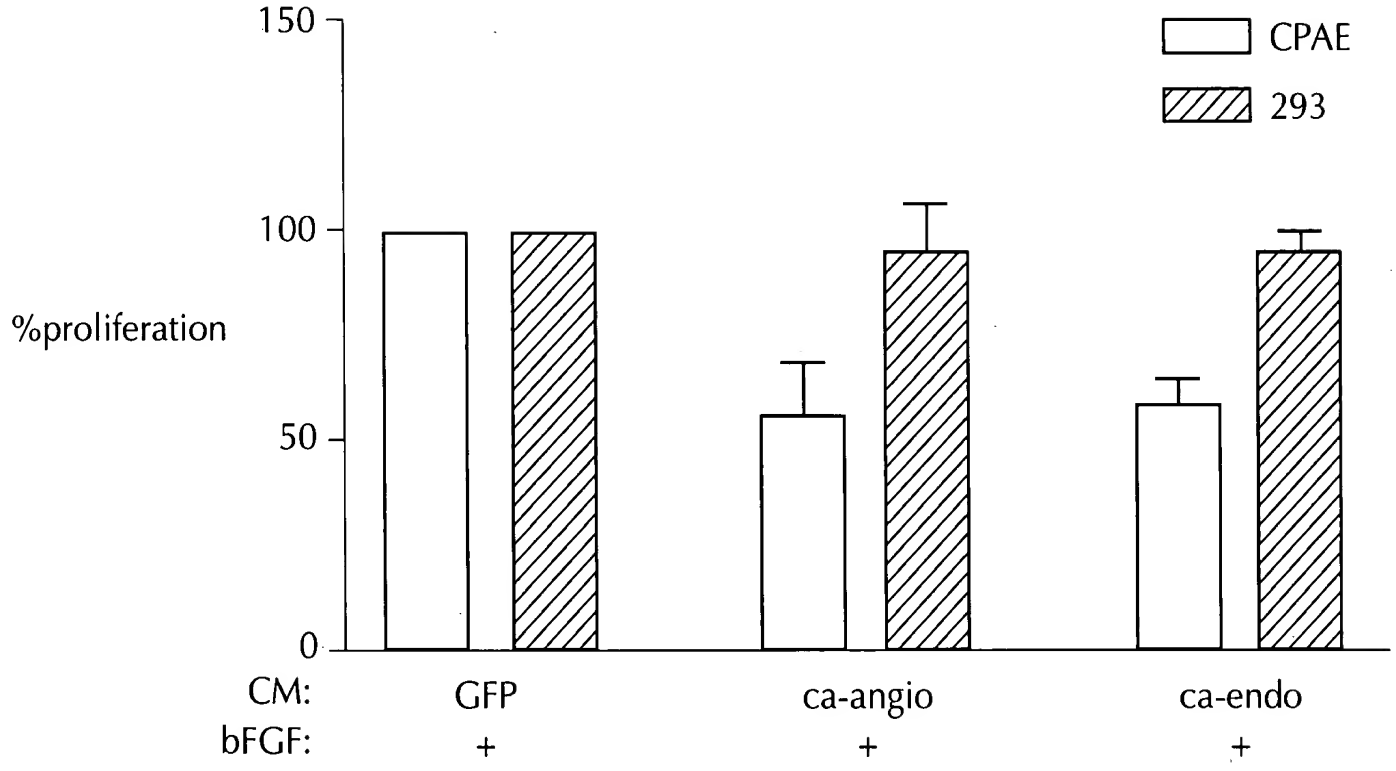


FIG. 10